## IN THE CLAIMS:

This listing of claims will replace all prior versions, and listing, of claims in the application.

## **Listing of the Claims:**

- 1. (Currently amended) A method of measurement of mitotic activity from histopathological specimen image data, characterised in that the method has comprising the steps of:
  - a) identifying pixels in the image data having luminances associated with mitotic figures;
  - b) selecting from among the identified pixels a reference pixel which is sufficiently close in position and luminance to another identified pixel to provide a reference colour;
  - c) locating pixels in the image data with luminances sufficiently close to that of the reference colour to indicate potentially mitotic figures;
  - d) incrementing image regions corresponding to potentially mitotic figures from the located pixels by adding pixels thereto, potential increments to image regions being implemented or rejected by according to whether or not their luminances are sufficiently close to respective image region luminances and sufficiently far from an image data background luminance;
  - e) selecting grown image regions on the basis of thresholds for image region area, compactness and width/height ratio; and
  - f) counting selected grown image regions as actually indicating mitotic figures on the basis of a threshold for number of such regions.
- 2. (Original) A method according to Claim 1 wherein the step of selecting grown image regions also involves thresholds for ratio of image region luminance to background luminance and area difference between areas derived by growing each image region with multiple thresholds.
- 3. (Original) A method according to Claim 2 wherein the thresholds for image region area,

compactness, width/height ratio, luminance and area difference are: 355 pixels < area < 1700 pixels, 0.17 < compactness < 0.77, width/height ratio < 2.7, luminance percentage < 44%, area difference < 23area/100.

- 4. (Currently amended) A method according to Claim 1 characterised in that wherein the step of counting selected grown image regions as actually indicating mitotic figures also involves thresholds for region area and luminance.
- 5. (Currently amended) A method according to Claim 1 characterised in that wherein successive potential increments to image regions are individual pixels each of which is an immediate row or column neighbour of an existing image region pixel.
- 6. (Currently amended) A method according to Claim 1 characterised in that wherein step
  b) is implemented with a reference pixel having a luminance differing by less than about
  8% compared to another identified pixel distant from it by not more than two percent of a smaller of two image dimensions.
- 7. (Currently amended) A method according to Claim 1 characterised in that wherein step a) includes white balancing and median filtering the image data prior to identifying pixels having luminances corresponding to mitotic figures.
- 8. (Currently amended) A method according to Claim 1 characterised in that wherein in step c) pixels are cued for acceptance or rejection as regards indicating mitotic figures by:
  - a) thresholding colour image data to remove pixels lacking intensities associated with mitotic figure imagery,
  - b) removal pixels not present in all colours, and
  - c) thresholding image region areas to remove those too small and too large to be potential mitotic figures.
- 9. (Currently amended) A method according to Claim 1 characterised in that wherein in step c) pixels are cued for acceptance or rejection as regards indicating mitotic figures by:

- a) segmenting to identify pixels with intensities associated with mitotic figure imagery,
- b) thresholding image region areas to remove those too small and too large to be potential mitotic figures,
- c) cluster analysis to determine whether or not a pixel's image region is in a sufficiently large cluster, and
- d) necrotic and hairy edge filtering.
- 10. (Currently amended) A method of measuring mitotic activity from histopathological specimen image data, characterised in that the method having the steps of:
  - a) measuring an intensity profile of an image region corresponding to a potentially mitotic figure, and
  - b) counting the image region as indicating a mitotic figure if its profile has a value greater than a prearranged threshold at a position in the profile having intensity associated with mitotic figure imagery.
- 11. (Currently amended) A method according to Claim 10 characterised in that it includes including counting the image region as indicating a mitotic figure if its profile has a first value not greater than the prearranged threshold at a position in the profile having intensity associated with mitotic figure imagery, a second value greater than a prearranged second threshold, a third value greater than a prearranged third threshold, and a minimum value less than a prearranged fourth threshold.
- 12. (Currently amended) A method according to Claim 11 characterised in that wherein the first value is at one end of the profile, the first and second values adjoin one another in the profile and the third value does not adjoin the second value.
- 13. (Currently amended) A method according to Claim 11 eharacterised in that wherein the image data comprise a first Principal Component obtained by Principal Component Analysis (PCA) of coloured image data.

- 14. (Currently amended) A method according to Claim 11 <del>characterised in that</del> wherein step a) includes preprocessing image data by:
  - a) decomposing the image data into overlapping sub-images,
  - b) applying PCA to the sub-images to derive a first Principal Component image,
  - c) thresholding the first Principal Component image to produce a binary image of blobs and background
  - d) rejecting blobs adjacent to or intersecting sub-image boundaries,
  - e) filling holes in blobs,
  - f) rejecting blobs too small to correspond to potential mitotic figures, and
  - g) reassembling the sub-images into a single image for image region profile measurement as aforesaid in step a).
- 15. (Currently amended) A method according to Claim 14 characterised in that wherein after step g) pixels are cued for acceptance or rejection as regards indicating mitotic figures by:
  - a) thresholding colour image data to remove pixels lacking intensities associated with mitotic figure imagery,
  - b) removal pixels not present in all colours, and
  - c) thresholding image region areas to remove those too small and too large to be potential mitotic figures.
- 16. (Currently amended) A method according to Claim 14 characterised in that wherein after step g) pixels are cued for acceptance or rejection as regards indicating mitotic figures by:
  - a) segmenting to identify pixels with intensities associated with mitotic figure imagery,
  - b) thresholding image region areas to remove those too small and too large to be potential mitotic figures,
  - c) cluster analysis to determine whether or not a pixel's image region is in a sufficiently large cluster, and
  - d) necrotic and hairy edge filtering.
- 17. (Currently amended) Computer apparatus for measuring mitotic activity from

histopathological specimen image data, <del>characterised in that it is</del> the apparatus being programmed to execute the steps of:

- a) identifying pixels in the image data having luminances associated with mitotic figures;
- b) selecting from among the identified pixels a reference pixel which is sufficiently close in position and luminance to another identified pixel to provide a reference colour;
- c) locating pixels in the image data with luminances sufficiently close to that of the reference colour to indicate potentially mitotic figures;
- d) incrementing image regions corresponding to potentially mitotic figures from the located pixels by adding pixels thereto, potential increments to image regions being implemented or rejected by according to whether or not their luminances are sufficiently close to respective image region luminances and sufficiently far from an image data background luminance;
- e) selecting grown image regions on the basis of thresholds for image region area, compactness and width/height ratio; and
- f) counting selected grown image regions as actually indicating mitotic figures on the basis of a threshold for number of such regions.
- 18. (Currently amended) Apparatus according to Claim 17 characterised in that it is programmed to execute the step of selecting grown image regions by also using thresholds for ratio of image region luminance to background luminance and area difference between areas derived by growing each image region with multiple thresholds.
- 19. (Currently amended) Apparatus according to Claim 18 eharacterised in that wherein the thresholds for image region area, compactness, width/height ratio, luminance and area difference are: 355 pixels < area < 1700 pixels, 0.17 < compactness < 0.77, width/height ratio < 2.7, luminance percentage < 44%, area difference < 23area/100.
- 20. (Currently amended) Apparatus according to Claim 17 <del>characterised in that it is</del> programmed to execute the step of counting selected grown image regions as actually

- indicating mitotic figures by also using thresholds for region area and luminance.
- 21. (Currently amended) Apparatus according to Claim 17 characterised in that wherein successive potential increments to image regions are individual pixels each of which is an immediate row or column neighbour of an existing image region pixel.
- 22. (Currently amended) Apparatus according to Claim 17 characterised in that it is programmed to execute step b) with a reference pixel having a luminance differing by less than 8% compared to another identified pixel distant from it by not more than two percent of a smaller of two image dimensions.
- 23. (Currently amended) Computer apparatus for measuring mitotic activity from histopathological specimen image data, characterised in that it is the apparatus being programmed to execute the steps of:
  - a) measuring an intensity profile of an image region corresponding to a potentially mitotic figure, and
  - b) counting the image region as indicating a mitotic figure if its profile has a value greater than a prearranged threshold at a position in the profile having intensity associated with mitotic figure imagery.
- 24. (Currently amended) Apparatus according to Claim 23 characterised in that it is also programmed to count an image region as indicating a mitotic figure if its profile has a first value not greater than the prearranged threshold at a position in the profile having intensity associated with mitotic figure imagery, a second value greater than a prearranged second threshold, a third value greater than a prearranged third threshold, and a minimum value less than a prearranged fourth threshold.
- 25. (Currently amended) Apparatus according to Claim 24 characterised in that wherein the first value is at one end of the profile, the first and second values adjoin one another in the profile and the third value does not adjoin the second value.
- 26. (Currently amended) Apparatus according to Claim 24 characterised in that wherein the

image data comprise a first Principal Component obtained by Principal Component Analysis (PCA) of coloured image data.

- 27. (Currently amended) A computer Computer program code for use in measuring mitotic activity from histopathological specimen image data, characterised in that the computer program code contains containing instructions to control a computer to implement the steps of:
  - a) identifying pixels in the image data having luminances associated with mitotic figures;
  - b) selecting from among the identified pixels a reference pixel which is sufficiently close in position and luminance to another identified pixel to provide a reference colour;
  - c) locating pixels in the image data with luminances sufficiently close to that of the reference colour to indicate potentially mitotic figures;
  - d) incrementing image regions corresponding to potentially mitotic figures from the located pixels by adding pixels thereto, potential increments to image regions being implemented or rejected by according to whether or not their luminances are sufficiently close to respective image region luminances and sufficiently far from an image data background luminance;
  - e) selecting grown image regions on the basis of thresholds for image region area, compactness and width/height ratio; and
  - f) counting selected grown image regions as actually indicating mitotic figures on the basis of a threshold for number of such regions.
- 28. (Currently amended) Computer program code according to Claim 27 eharacterised in that its incorporating instructions to provide for implementing the step of selecting grown image regions by also using thresholds for ratio of image region luminance to background luminance and area difference between areas derived by growing each image region with multiple thresholds.
- 29. (Currently amended) Computer program code according to Claim 28 characterised in that

wherein the thresholds for image region area, compactness, width/height ratio, luminance and area difference are: 355 pixels < area < 1700 pixels, 0.17 < compactness < 0.77, width/height ratio < 2.7, luminance percentage < 44%, area difference < 23area/100.

- 30. (Currently amended) A computer Computer program code according to Claim 27 characterised in that its incorporating instructions to provide for implementing the step of counting selected grown image regions as actually indicating mitotic figures using also thresholds for region area and luminance.
- 31. (Currently amended) A-computer Computer program code for use in measuring mitotic activity from histopathological specimen image data, characterised in that its incorporating instructions to provide for implementing the steps of:
  - a) measuring an intensity profile of an image region corresponding to a potentially mitotic figure, and
  - b) counting the image region as indicating a mitotic figure if its profile has a value greater than a prearranged threshold at a position in the profile having intensity associated with mitotic figure imagery.
- 32. (Currently amended) A computer program according to Claim 31 characterised in that its incorporating instructions to provide for counting the image region as indicating a mitotic figure if its profile has a first value not greater than the prearranged threshold at a position in the profile having intensity associated with mitotic figure imagery, a second value greater than a prearranged second threshold, a third value greater than a prearranged third threshold, and a minimum value less than a prearranged fourth threshold.
- 33. (Currently amended) A computer program according to Claim 32 characterised in that wherein the first value is at one end of the profile, the first and second values adjoin one another in the profile and the third value does not adjoin the second value.
- 34. (Currently amended) A computer program according to Claim 32 <del>characterised in that its</del> incorporating instructions to provide for step a) to include preprocessing image data by:

- a) decomposing the image data into overlapping sub-images,
- b) applying PCA to the sub-images to derive a first Principal Component image,
- c) thresholding the first Principal Component image to produce a binary image of blobs and background
- d) rejecting blobs adjacent to or intersecting sub-image boundaries,
- e) filling holes in blobs,
- f) rejecting blobs too small to correspond to potential mitotic figures, and
- g) reassembling the sub-images into a single image for image region profile measurement as aforesaid in step a).